f 7

SEQUENCE LISTING

<110> Croze, Ed
     Vogel, David
     Russell-Harde, Dean

<120> THE USE OF AN INTERFERON RECEPTOR POLYPEPTIDE
     CHAIN TO ENHANCE THE EFFECTS OF INTERFERONS

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<170> PatentIn Ver. 2.1

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Asp Lys Trp Leu Glu Gly Met Glu Ser Tyr Val Val Ala Ile Val Ile
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Val His Arg Gly Asp Leu Thr Val Cys Arg Cys Ser Asp Tyr Ile Val
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Pro Ala Asn Ala Pro Leu Glu Pro Pro Glu Phe Glu Ile Val Gly Phe
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Leu Lys Cys Ile Val Leu Gln Pro Gly Gln Glu Ser Gly Leu Ser Glu
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Val Ser Thr Ile Val Met Leu Lys Arg Ile Gly Tyr Ile Cys Leu Lys
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Asp Asn Leu Pro Asn Val Leu Asn Phe Arg His Phe Leu Thr Trp Ile
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Ile Pro Glu Arg Ser Pro Ser Glu Ala Ile Asp Arg Leu Glu Ile Ile
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Pro Thr Asn Lys Lys Lys Arg Leu Trp Asn Tyr Asp Tyr Glu Asp Gly
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Arg Lys Ser Val Leu Glu Asp Ser Phe Pro Arg Glu Asp Asn Ser Ser
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Ser Asp Leu Arg Ile Ala Gly Gly Asp Arg Thr Gln Pro Pro Leu Pro
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Asp Tyr Thr Asp Glu Ser Cys Thr Phe Lys Ile Ser Leu Arg Asn Phe
         35                  40                  45

Arg Ser Ile Leu Ser Trp Glu Leu Lys Asn His Ser Ile Val Pro Thr
     50                  55                  60

His Tyr Thr Leu Leu Tyr Thr Ile Met Ser Lys Pro Glu Asp Leu Lys
 65                  70                  75                  80

Val Val Lys Asn Cys Ala Asn Thr Thr Arg Ser Phe Cys Asp Leu Thr
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Asp Glu Trp Arg Ser Thr His Glu Ala Tyr Val Thr Val Leu Glu Gly
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Phe Ser Gly Asn Thr Thr Leu Phe Ser Cys Ser His Asn Phe Trp Leu
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Ala Ile Asp Met Ser Phe Glu Pro Pro Glu Phe Glu Ile Val Gly Phe
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Glu Leu Gln Phe Asp Leu Ser Leu Val Ile Glu Glu Gln Ser Glu Gly
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Ile Val Lys Lys His Lys Pro Glu Ile Lys Gly Asn Met Ser Gly Asn
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Phe Thr Tyr Ile Ile Asp Lys Leu Ile Pro Asn Thr Asn Tyr Cys Val
         195                 200                 205

Ser Val Tyr Leu Glu His Ser Asp Glu Gln Ala Val Ile Lys Ser Pro
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Leu Lys Cys Thr Leu Leu Pro Pro Gly Gln Glu Ser Glu Ser Ala Glu
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Ser Ala Lys Ile Gly Gly Ile Ile Thr Val Phe Leu Ile Ala Leu Val
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Leu Thr Ser Thr Ile Val Thr Leu Lys Trp Ile Gly Tyr Ile Cys Leu
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Pro Phe Pro Asn Leu Pro Pro Leu Glu Ala Met Asp Met Val Glu Val
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Ile Tyr Ile Asn Arg Lys Lys Lys Val Trp Asp Tyr Asn Tyr Asp Asp
305   310   315   320

Glu Ser Asp Ser Asp Thr Glu Ala Ala Pro Arg Thr Ser Gly Gly Gly
            325   330   335

Tyr Thr Met His Gly Leu Thr Val Arg Pro Leu Gly Gln Ala Ser Ala
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Thr Ser Thr Glu Ser Gln Leu Ile Asp Pro Glu Ser Glu Glu Glu Pro
        355   360   365

Asp Leu Pro Glu Val Asp Val Glu Leu Pro Thr Met Pro Lys Asp Ser
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Pro Gln Gln Leu Glu Leu Leu Ser Gly Pro Cys Glu Arg Arg Lys Ser
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Pro Leu Gln Asp Pro Phe Pro Glu Glu Asp Tyr Ser Ser Thr Glu Gly
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Ser Gly Gly Arg Ile Thr Phe Asn Val Asp Leu Asn Ser Val Phe Leu
            420   425   430

Arg Val Leu Asp Asp Glu Asp Ser Asp Asp Leu Glu Ala Pro Leu Met
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Leu Ser Ser His Leu Glu Glu Met Val Asp Pro Glu Asp Pro Asp Asn
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Val Gln Ser Asn His Leu Leu Ala Ser Gly Glu Gly Thr Gln Pro Thr
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Phe Pro Ser Pro Ser Ser Glu Gly Leu Trp Ser Glu Asp Ala Pro Ser
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